PAGE: 1 Sohn Servi

RAW SEQUENCE LISTING PATENT APPLICATION US/09/052,855A

DATE: 08/23/98 TIME: 14:29:12

INPUT SET: S28208.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

	,1				SEQUENC	E LISTING	,
	2		- 7				r.
	3	(1)	General	Information			ENTERED
	4 5		/i)	ADDITCANT. B	ILLING-MEDEL,	DATRICTA	LAED
	6		(-)	COHEN,	= = = = = = = = = = = = = = = = = = = =	TATRICIA	
	7			-	S, TRACEY L.		•
	8				N, PAULA N.		
	9			GORDON,			
	10				S, EDWARD N.		
	11			HAYDEN,	MARK		
	12			HODGES,	STEVEN C.		
	13			KLASS,	MICHAEL R.		
	14				VIL, JON D.		
	15				-RAPP, LISA		
>	16				, JOHN C.		
>	17			STROUPE	, STEPHEN D.		
	18						
	19		(11)			REAGENTS AND M	
	20 21				CTING DISEASE	S OF THE GAST	ROINTESTINAL
	21			TRACT			
	23		(1111)	MIMPED OF C	POHENCEC. 20		
	24		(111)	NUMBER OF S	EQUENCES: 30		
	25		/ i v)	CORRESPONDEN	CE ADDRESS.		
	26		• •		Abbott Labora	tories	
	27		V /		Abbott Park		
	28			CITY: Abbot			
	29		• •	STATE: IL			
	30			COUNTRY: US.	A		
	31	•	(F)	ZIP: 60064-	3500		
	32		•				
	33		(V) C	OMPUTER READ	ABLE FORM:		
	34		(A)	MEDIUM TYPE	: Diskette		
	35		• •		BM Compatible	!	
	36		, ,	OPERATING S			
	37		(D)	SOFTWARE: F	astSEQ for Wi	ndows Version	2.0
	38						
	39		•		ICATION DATA:		
	40				NUMBER: 09/0		
	41				: 31-MAR-1998		
	42		(C)	CLASSIFICAT	TON:		
	43 44		/ 2 - 1	מסדסם אחתי די	באשדרו וויאשא.		
	44 45		•	PRIOR APPLI		28 489	
	45 46				NUMBER: 08/8 : 31-MAR-1997	·	
	40		(B)	EILING DATE	. JI-MAK-177/		

RAW SEQUENCE LISTING PATENT APPLICATION US/09/052,855A

DATE: 08/23/98 TIME: 14:29:13

47				
48	(viii) ATTORNEY/AGENT INFORMATION:			
49	(A) NAME: Becker, Cheryl L.			
50	(B) REGISTRATION NUMBER: 35,441			
51	(C) REFERENCE/DOCKET NUMBER: 6064.US.Pl			
52				
53	(ix) TELECOMMUNICATION INFORMATION:			
54	(A) TELEPHONE: 847/935-1729			
55	(B) TELEFAX: 847/938-2623			
56	(C) TELEX:			
57	, ,			
58	(2) INFORMATION FOR SEQ ID NO:1:			
59				
60	(i) SEQUENCE CHARACTERISTICS:			
61	(A) LENGTH: 273 base pairs			
62	(B) TYPE: nucleic acid			
63	(C) STRANDEDNESS: single			
64	(D) TOPOLOGY: linear			
65	(-)			
66	(ix) FEATURE:			
67	(A) NAME/KEY: base_polymorphism			
68	(B) LOCATION: 3			
69	(D) OTHER INFORMATION: /note= " N' represents an A or G or			
70	T or C polymorphism at this position"			
71				
72	(ix) FEATURE:			
73	(A) NAME/KEY: base polymorphism			
74	(B) LOCATION: 55			
75	(D) OTHER INFORMATION: /note= " N' represents an A or G or			
76	T or C polymorphism at this position"			
77	Fallman Francisco			
78	(ix) FEATURE:			
79	(A) NAME/KEY: base polymorphism			
80	(B) LOCATION: 62			
81	(D) OTHER INFORMATION: /note= " N' represents an A or G or			
82	T or C polymorphism at this position"			
83	2 02 0 902,29			
84	(ix) FEATURE:			
85	(A) NAME/KEY: base polymorphism			
86	(B) LOCATION: 189			
87	(D) OTHER INFORMATION: /note= " N' represents an A or G or			
88	T or C polymorphism at this position"			
89	Tot o potymorphism at this postation			
90	(ix) FEATURE:			
91	(A) NAME/KEY: base_polymorphism			
92	(B) LOCATION: 201			
93	(D) OTHER INFORMATION: /note= " N' represents an A or G or			
94	T or C polymorphism at this position"			
95	1 of o polymorphism at this position			
96	(ix) FEATURE:			
97	(A) NAME/KEY: base_polymorphism			
98	(B) LOCATION: 204			
99	(D) OTHER INFORMATION: /note= " N' represents an A or G or			
,,	(P) OTHER THEOREMITOR: \HOCE H TEDIESEHES ON WOL O			

RAW SEQUENCE LISTING PATENT APPLICATION US/09/052,855A

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	INPUT SET: S28208.raw
100	T or C polymorphism at this position"
101	
102	(ix) FEATURE:
103	(A) NAME/KEY: base polymorphism
104	(B) LOCATION: 206
105	(D) OTHER INFORMATION: /note= " N' represents an A or G or
106	T or C polymorphism at this position"
107	Tot e polymorphism at this position
108	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
109	(XI) SECONDE DESCRIPTION. SEC ID NO. 1.
110	GGNCAGAGCC TGCGCAGGGC AGGAGCAGCT GGCCCACTGG CGGCCCGCAA CACTNCGTCT 60
	TNACCCTCTG GGCCCACTGC ATCTAGAGGA GGGCCGTCTG TGAGGCCACT ACCCCTCCAG 120
111	•
112	
113	CTACGGCANC TGCTGGACCA NCTNGNCTTT TCCATCGAAG CAGGGAAGTG GGAGCCTTGA 240
114	GCCCTTGGGT GGAAGCTTGA CCCCAAGCCA CTT 273
115	
116	(2) INFORMATION FOR SEQ ID NO:2:
117	
118	(i) SEQUENCE CHARACTERISTICS:
119	(A) LENGTH: 250 base pairs
120	(B) TYPE: nucleic acid
121	(C) STRANDEDNESS: single
122	(D) TOPOLOGY: linear
123	
124	(ix) FEATURE:
125	(A) NAME/KEY: base_polymorphism
126	(B) LOCATION: 69
127	(D) OTHER INFORMATION: /note= " N' represents an A or G or
128	T or C polymorphism at this position"
129	• • •
130	(ix) FEATURE:
131	(A) NAME/KEY: base polymorphism
132	(B) LOCATION: 97
133	(D) OTHER INFORMATION: /note= " N' represents an A or G or
134	T or C polymorphism at this position"
135	to the second se
136	(ix) FEATURE:
137	(A) NAME/KEY: base polymorphism
138	(B) LOCATION: 140
139	(D) OTHER INFORMATION: /note= " N' represents an A or G or
140	T or C polymorphism at this position"
	1 of c polymorphism at this position
141	(in) BRANCE.
142	(ix) FEATURE:
143	(A) NAME/KEY: base_polymorphism
144	(B) LOCATION: 223
145	(D) OTHER INFORMATION: /note= " N' represents an A or G or
146	T or C polymorphism at this position"
147	
148	
149	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
150	
151	AGAGCCTGCG CAGGGCAGGA GCAGCTGGCC CACTGGCGGC CCGCAACACT CCGTCTCACC 60
152	CTCTGGGCNC ACTGCATCTA GAGGAGGGCC GTCTGTNAGG CCACTACCCC TCCAGCAACT 120

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153 154	INPUT SET: S28208.raw GGGAGGTGGG ACTGTCAGAN GCTGGCCCAG GGTGGTGGTC AGCTGGGTCA GGGACCTACG 180 GCACCTGCTG GACCACCTCG CCTTCTCCAT CGAAGCAGGG AANTGGGAGC CTCGAGCCCT 240
155	CGGGTGGAAG 250
156	
157	(2) INFORMATION FOR SEQ ID NO:3:
158	
159	(i) SEQUENCE CHARACTERISTICS:
160	(A) LENGTH: 250 base pairs
161	(B) TYPE: nucleic acid
162	(C) STRANDEDNESS: single
163	(D) TOPOLOGY: linear
164	
165	(ix) FEATURE:
166	(A) NAME/KEY: base_polymorphism
167	(B) LOCATION: 68
168	(D) OTHER INFORMATION: /note= " N' represents an A or G or
169	T or C polymorphism at this position"
170	
171	(ix) FEATURE:
172	(A) NAME/KEY: base polymorphism
173	(B) LOCATION: 232
174	(D) OTHER INFORMATION: /note= " N' represents an A or G or
175	T or C polymorphism at this position"
176	
177	(ix) FEATURE:
178	(A) NAME/KEY: base polymorphism
179	(B) LOCATION: 233
180	(D) OTHER INFORMATION: /note= " N' represents an A or G or
181	T or C polymorphism at this position"
182	o or o god, morphism and annual grand and a
183	
184	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
185	()
186	TGGCGGCCCG CAACACTCCG TCTCACCCTC TGGGCCCACT GCATCTAGAG GAGGGCCGTC 60
187	TGTGAGGNCA CTACCCCTCC AGCAACTGGG AGGTGGGACT GTCAGAATCT GGCCCAGGGT 120
188	GGTGGTCAGC TGGGTCAGGG ACCTACGGCA CCTGCTGGAC CACCTCGCCT TCTCCATCGA 180
189	AGCAGGGAAG TGGGAGCCTC GAGCCCTCGG GTGGAAGCTG ACCCCAAGCC ANNCTTCACC 240
190	TGGACAGGAT 250
191	
192	(2) INFORMATION FOR SEQ ID NO:4:
193	1.2/
194	(i) SEQUENCE CHARACTERISTICS:
195	(A) LENGTH: 233 base pairs
196	(B) TYPE: nucleic acid
197	(C) STRANDEDNESS: single
198	(D) TOPOLOGY: linear
199	12/ 10102011 11:001
200	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
201	(vr) ppsopued pppovirizion, pps in no.1.
201	CCCTCTGGGC CCACTGCATC TAGAGGAGGG CCGTCTGTGA GGCCACTACC CCTCCAGCAA 60
202	CTGGGAGGTG GGACTGTCAG AAGCTGGCCC AGGGTGGTGG TCAGCTGGGT CAGGGACCTA 120
203	CGGCACCTGC TGGACCACCT CGCCTTCTCC ATCGAAGCAG GGAAGTGGGA GCCTCGAGCC 180
204	CTCGGGTGGA AGCTGACCC AAGCCACCCT TCACCTGGAC AGGATGAGAG TGT 233
203	CICOGGIGGA AGCIGACCCC AAGCCACCCI ICACCIGGAC AGGAIGAGAG IGI 233

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206	
207	(2) INFORMATION FOR SEQ ID NO:5:
208	
209	(i) SEQUENCE CHARACTERISTICS:
210	(A) LENGTH: 287 base pairs
211	(B) TYPE: nucleic acid
212	(C) STRANDEDNESS: single
213	(D) TOPOLOGY: linear
214	. (5) 10102011 121011
215	(ix) FEATURE:
216	(A) NAME/KEY: base polymorphism
217	(B) LOCATION: 17
218	(D) OTHER INFORMATION: /note= " N' represents an A or G or
219	T or C polymorphism at this position"
220	1 of C polymorphism at this position
	(iv) PRAMUDD.
221	(ix) FEATURE:
222	(A) NAME/KEY: base_polymorphism
223	(B) LOCATION: 60
224	(D) OTHER INFORMATION: /note= " N' represents an A or G or
225	T or C polymorphism at this position"
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227	(ix) FEATURE:
228	(A) NAME/KEY: base_polymorphism
229	(B) LOCATION: 193
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231	T or C polymorphism at this position"
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233	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
234	,
235	CACGAGGGCC GTCTGTNAGG CCACTACCCC TCCAGCAACT GGGAGGTGGG ACTGTCAGAN 60
236	GCTGGCCCAG GGTGGTGGTC AGCTGGGTCA GGGACCTACG GCACCTGCTG GACCACCTCG 120
237	CCTTCTCCAT CGAAGCAGGG AAGTGGGAGC CTCGAGCCCT CGGGTGGAAG CTGACCCCAA 180
238	GCCACCCTTC ACNTGGACAG GATGAGAGTG TCAGGTGTGC TTCGCCTCCT GGCCCTCATC 240
239	TTTGCCATAG TCACGACATG GATGTTTATT CGAAGCTACA TGAGCTT 287
240	
241	(2) INFORMATION FOR SEQ ID NO:6:
242	
243	(i) SEQUENCE CHARACTERISTICS:
244	(A) LENGTH: 263 base pairs
245	(B) TYPE: nucleic acid
246	(C) STRANDEDNESS: single
247	(D) TOPOLOGY: linear
248	
249	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
250	· · · · · · · · · · · · · · · · · · ·
251	GATGTTTATT CGAAGCTACA TGAGCTTCAG CATGAAAACC ATCCGTCTGC CACGCTGGCT 60
252	GGCCTCGCCC ACCAAGGAGA TCCAGGTTAA AAAGTACAAG TGTGGCCTCA TCAAGCCCTG 120
253	CCCAGCCAAC TACTTTGCGT TTAAAATCTG CAGTGGGGCC GCCAACGTCG TGGGCCCTAC 180
254	TATGTGCTTT GAAGACCGCA TGATCATGAG TCCTGTGAAA AACAATGTGG GCAGAGGCCT 240
255	AAACATCGCC CTGGTGAATG GAA 263
256	
257	(2) INFORMATION FOR SEQ ID NO:7:
258	(2) INTORMATION FOR DBQ ID NO. /.
200	

SEQUENCE VERIFICATION REPORT PATENT APPLICATION *US/09/052,855A*

DATE: 08/23/98 TIME: 14:29:17

Line	Error	Original Text
16 17	Response Exceeds Line Limitations Response Exceeds Line Limitations	RUSSELL, JOHN C. STROUPE, STEPHEN D.